

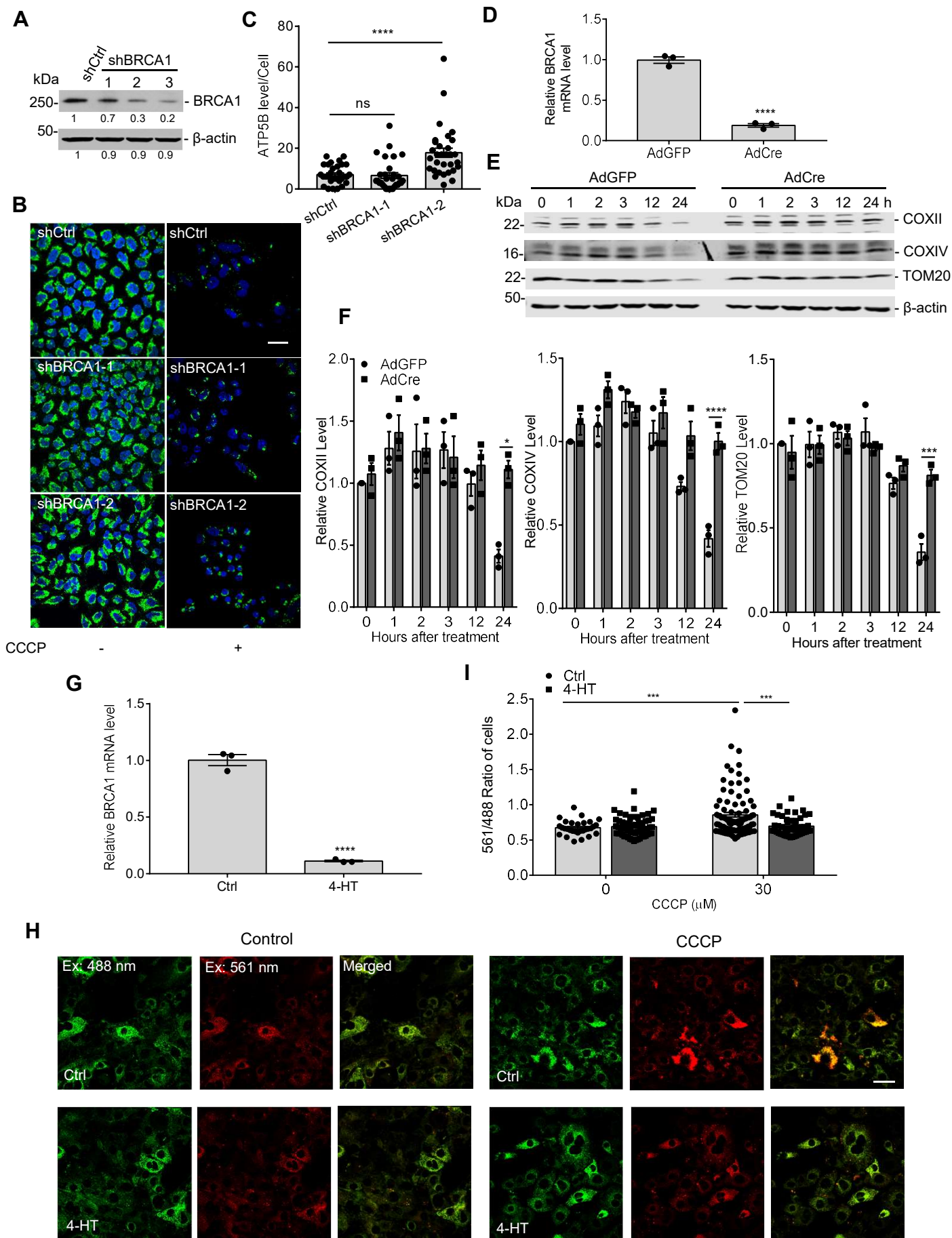


## Supporting Information

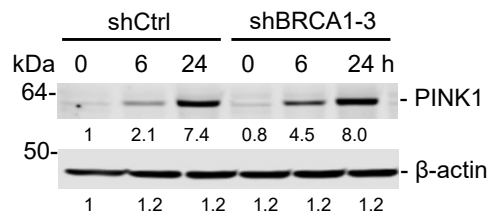
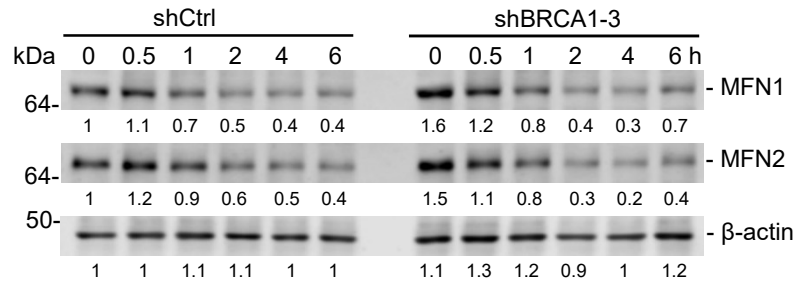
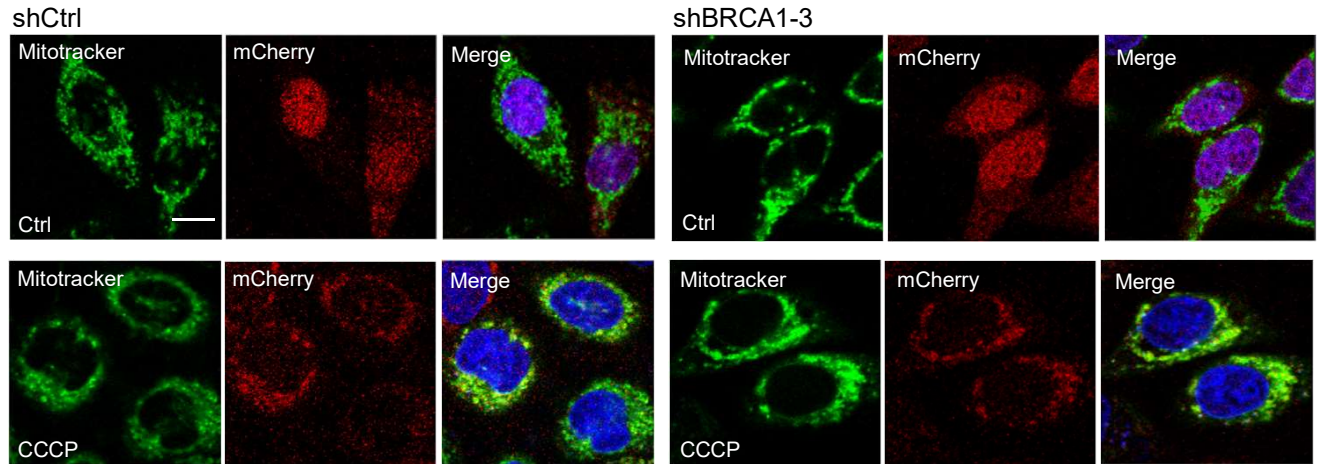
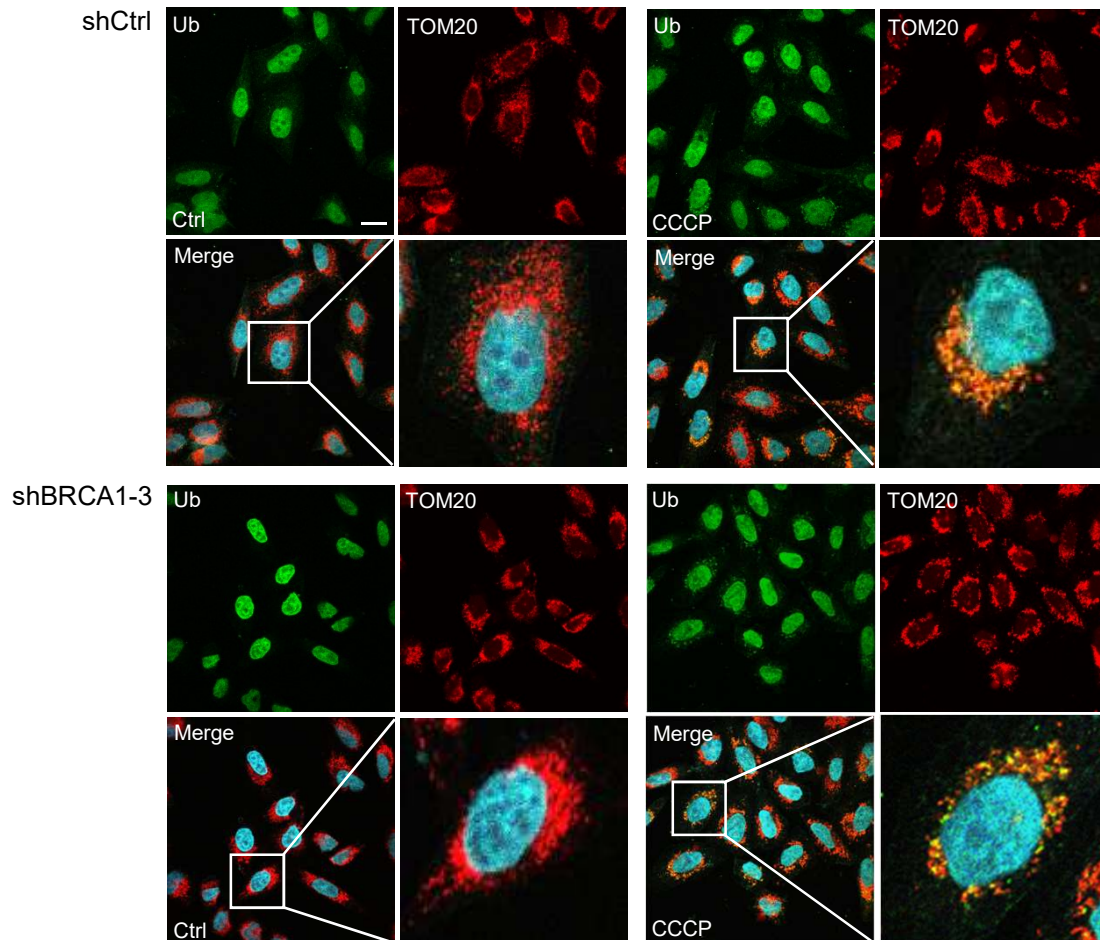
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**BRCA1 Deficiency Impairs Mitophagy and Promotes  
Inflammasome Activation and Mammary Tumor Metastasis**

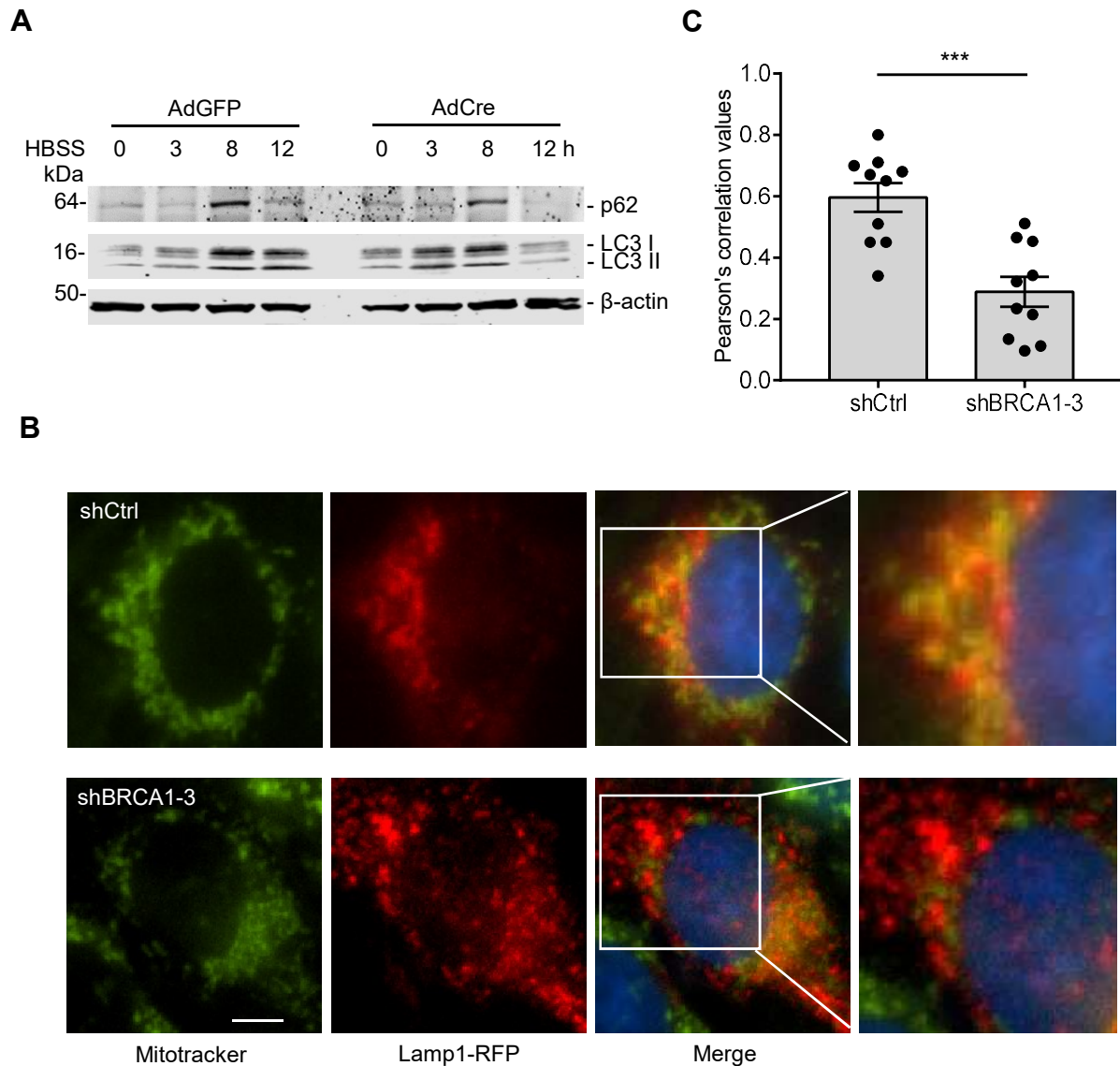
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**Figure S1. Loss of BRCA1 impairs stress-induced mitophagy.** **A)** Immunoblot analysis of BRCA1 in Hela-mCherryParkin with control shRNA or different BRCA1 shRNA. **B)** Analysis of mitophagy activities in shCtrl and shBRCA1 Hela-mCherryParkin under CCCP treatment by clearance of ATP5B. Scale bar, 20  $\mu$ m. **C)** Quantification for ATP5B levels after CCCP treatment (more than 30 cells were counted per group). **D)** The mRNA levels of BRCA1 in *Brca1<sup>flox/flox</sup>* MEFs infected with AdGFP or AdCre. **E)** Immunoblot analysis of COXII, COXIV and TOM20 in *Brca1<sup>flox/flox</sup>* MEFs infected with AdGFP or AdCre treated with CCCP. **F)** Quantification of COXII, COXIV and TOM20 levels in (D), which normalized by  $\beta$ -actin level (n = 3 per group). **G)** The mRNA levels of BRCA1 in *Brca1<sup>flox/flox</sup>;Tam-Cre* MEFs with or without 4-HT treatment. **H, I)** Mitophagy activity indicated by mt-mKeima in *Brca1<sup>flox/flox</sup>;Tam-Cre* MEFs. (H) Dual-excitation imaging (488/561 nm) of mt-mKeima. Scale bar, 50  $\mu$ m. (I) Quantification of fluorescence intensity from (H) (more than 50 cells were counted per group). A high ratio (561/488) of puncta indicates a low pH area. The cells were treated with or not CCCP (30  $\mu$ M). Data represent the mean  $\pm$  SEM and are representative of three independent experiments. Significant differences were determined by one-way (C) or two-way (F, I) ANOVA with Tukey multiple comparison testing or unpaired two-tailed *t* test (D and G). \**p* < 0.05, \*\*\**p* < 0.001, \*\*\*\**p* < 0.0001.

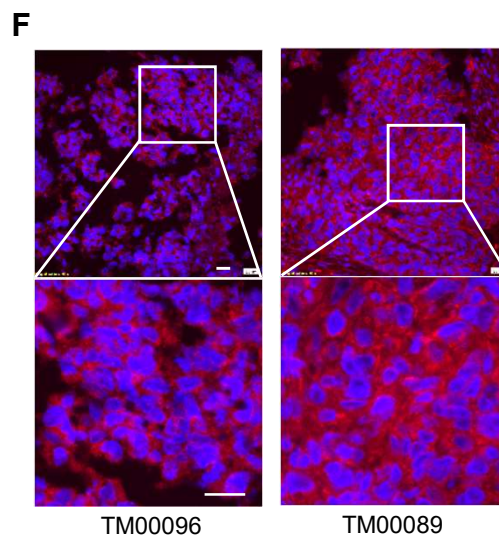
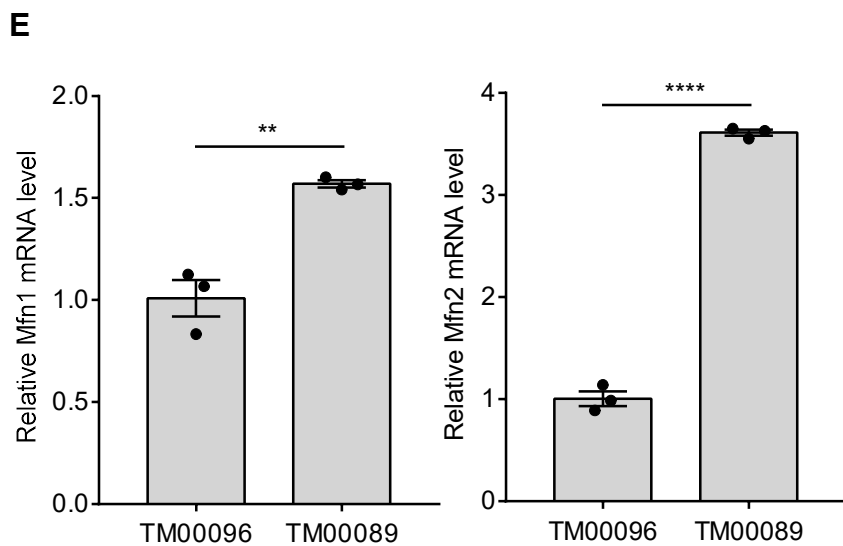
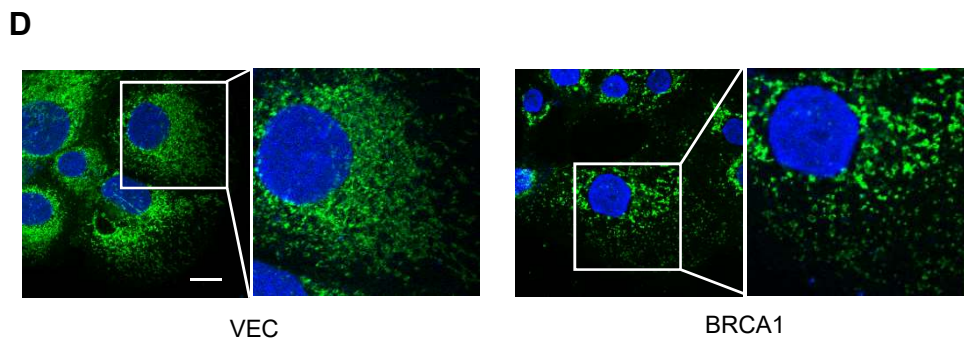
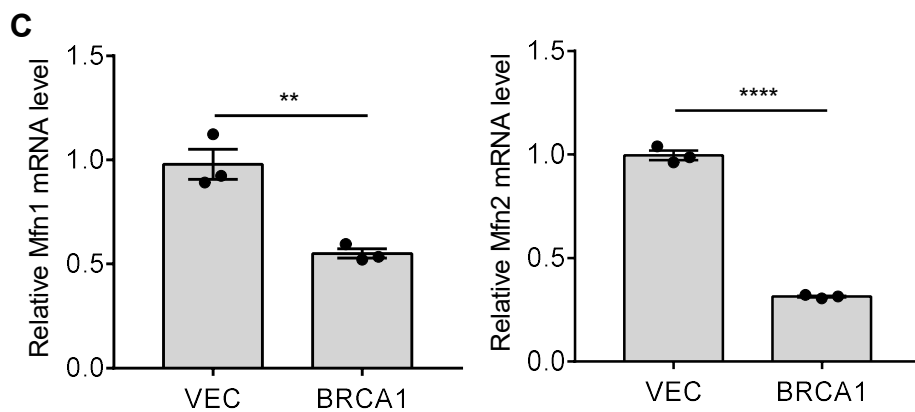
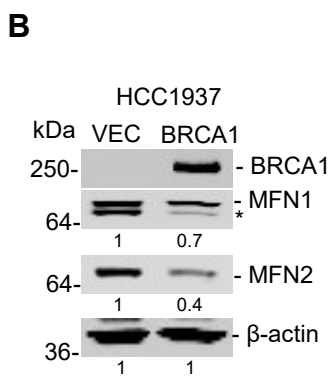
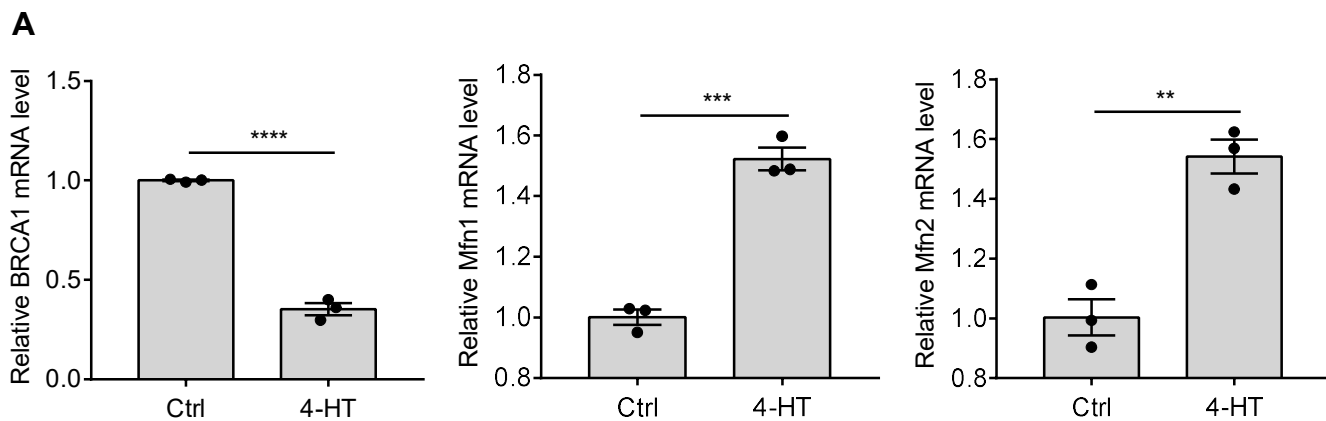
**A****D****B****C**

**Figure S2. BRCA1 deficiency has no effect on CCCP induced PINK1/Parkin pathway activation.** **A)** Immunoblot analysis of PINK1 in shCtrl and shBRCA1 HeLa cells after CCCP (10  $\mu$ M) treatment for indicated time points. **B)** shCtrl and shBRCA1 HeLa-mCherryParkin were treated by CCCP for 1 h, and then stained by MitoTracker Green for labeling mitochondria. Hoechst33258, DNA-binding dye. Scale bar, 10  $\mu$ m. **C)** The ubiquitin level on mitochondria in shCtrl and shBRCA1 HeLa-HA-Parkin with or without CCCP (10  $\mu$ M) treatment for 3 h, as measured by immunostaining for ubiquitin (Ub). Mitochondria were labeled by TOM20 immunostaining. Scale bar, 10  $\mu$ m. **D)** Immunoblot analysis of MFN1 and MFN2 levels in HeLa-HA-Parkin shCtrl and shBRCA1 cells under CCCP treatment for indicated time points. Data are representative of at least two independent experiments.



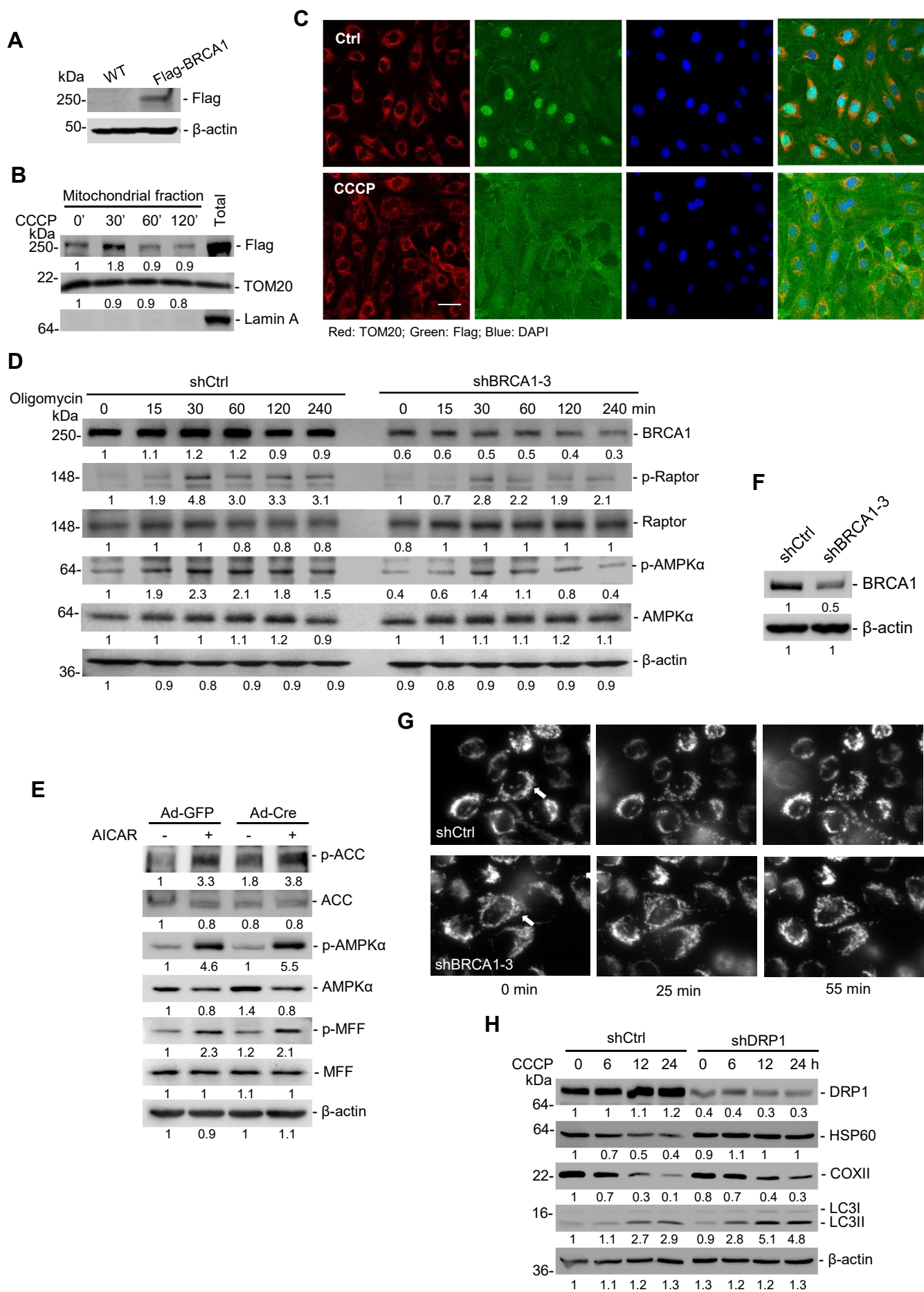
**Figure S3. BRCA1 is required for mitophagy, but not for general autophagy.** **A)** Immunoblot analysis of p62 and LC3 in *Brca1<sup>flox/flox</sup>* MEFs infected with AdGFP or AdCre under starvation (HBSS treatment) for indicated times. **B)** The co-localization of lysosomes with mitochondria in HeLa-HA-Parkin shCtrl and shBRCA1 cells under CCCP (10  $\mu$ M) treatment for 6 h. Lysosomes and mitochondria were marked with LAMP1-RFP and MitoTracker Green, respectively. Scale bar, 5  $\mu$ m. **C)** Pearson's coefficient is shown as the quantification of lysosomes co-localized with mitochondria per cell in (B) (10 fields were counted per group). Data represent the mean  $\pm$  SEM and are representative of at least two independent experiments. Significant differences were determined by unpaired two-tailed *t* test (C) \*\*\* $p < 0.001$ .



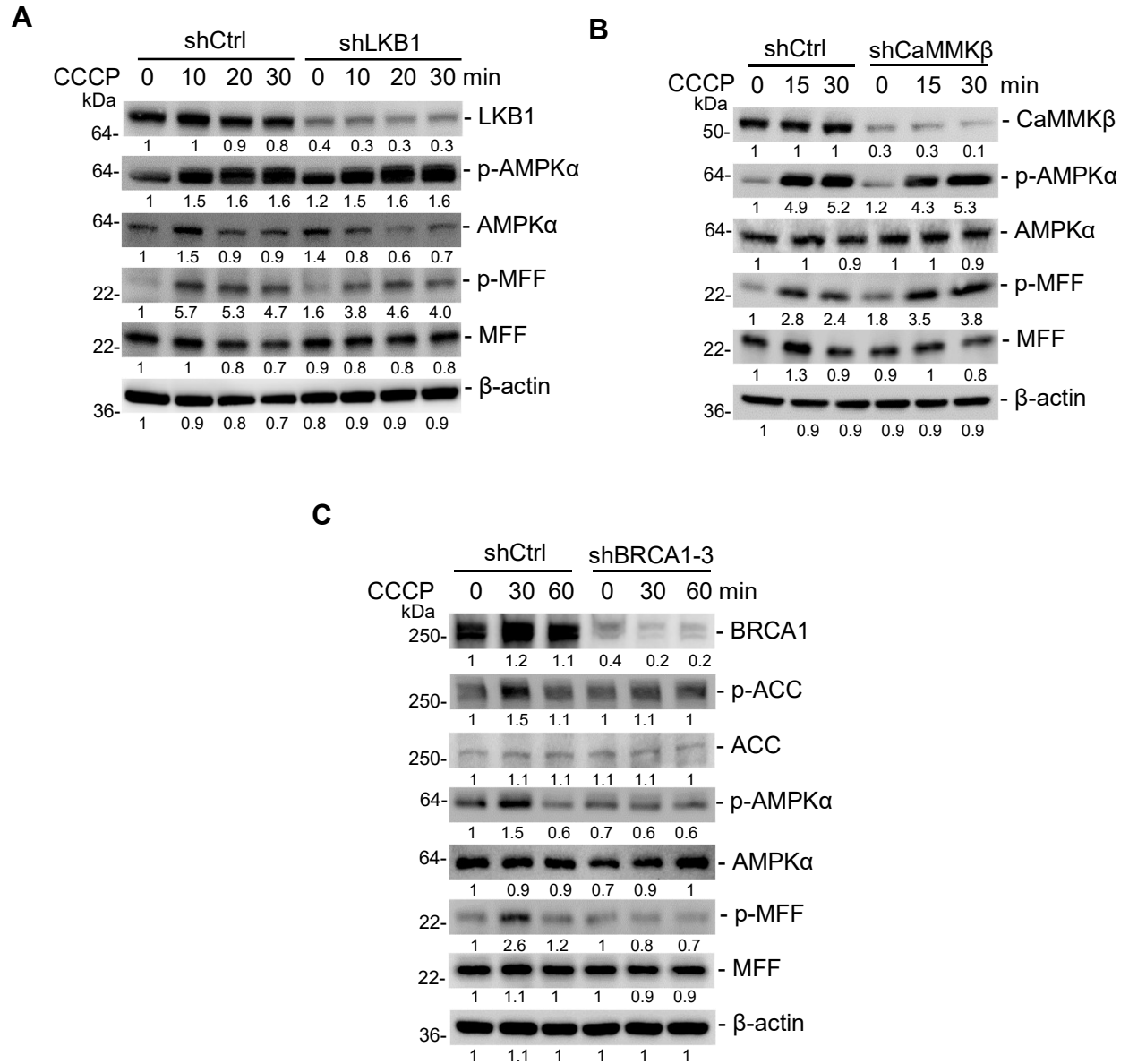


**Figure S4. BRCA1 regulates MFN1/2 mediated-mitochondrial fusion.** **A)** The mRNA levels of BRCA1, Mfn1 and Mfn2 in *Brca1<sup>flox/flox</sup>; Tam-Cre* MEFs with or without 4-HT (5  $\mu$ M) treatment, as measured by real-time PCR (n = 3 per group). **B)** Immunoblot analysis of BRCA1, MFN1 and MFN2 in HCC1937 cells stably transfected with empty vector (VEC) or BRCA1 expressing vector (BRCA1). Asterisk (\*) indicates MFN2 band. **C)** The mRNA levels of Mfn1 and Mfn2 in VEC and BRCA1 HCC1937 cells measured by real-time PCR (n = 3 per group). **D)** Mitochondrial morphology of VEC and BRCA1 HCC1937 cells stained by MitoTracker Green. Scale bar, 10  $\mu$ m. **E)** The mRNA levels of Mfn1 and Mfn2 in breast cancer PDX models (TM00096, *BRCA1* WT; TM00089, *BRCA1* MT) cells measured by real-time PCR (n = 3 per group). **F)** Representative images of immunostained MFN1 in breast cancer PDX models. Scale bar, 20  $\mu$ m. Data represent the mean  $\pm$  SEM and are representative of at least two independent experiments. Significant differences were determined by unpaired two-tailed *t* test. \*\**p* < 0.01, \*\*\**p* < 0.001, \*\*\*\**p* < 0.0001.

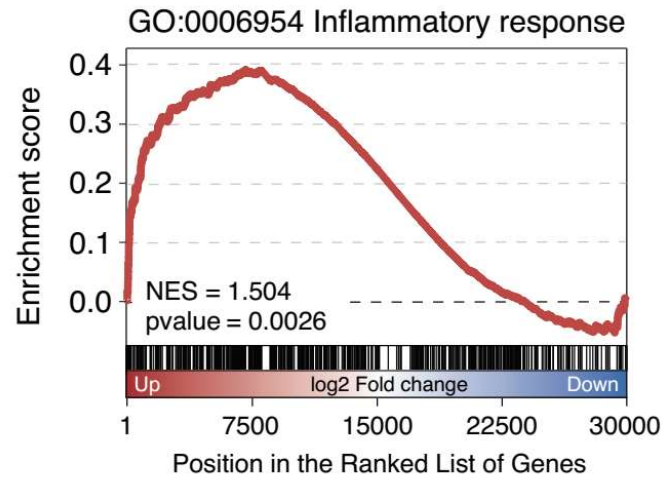




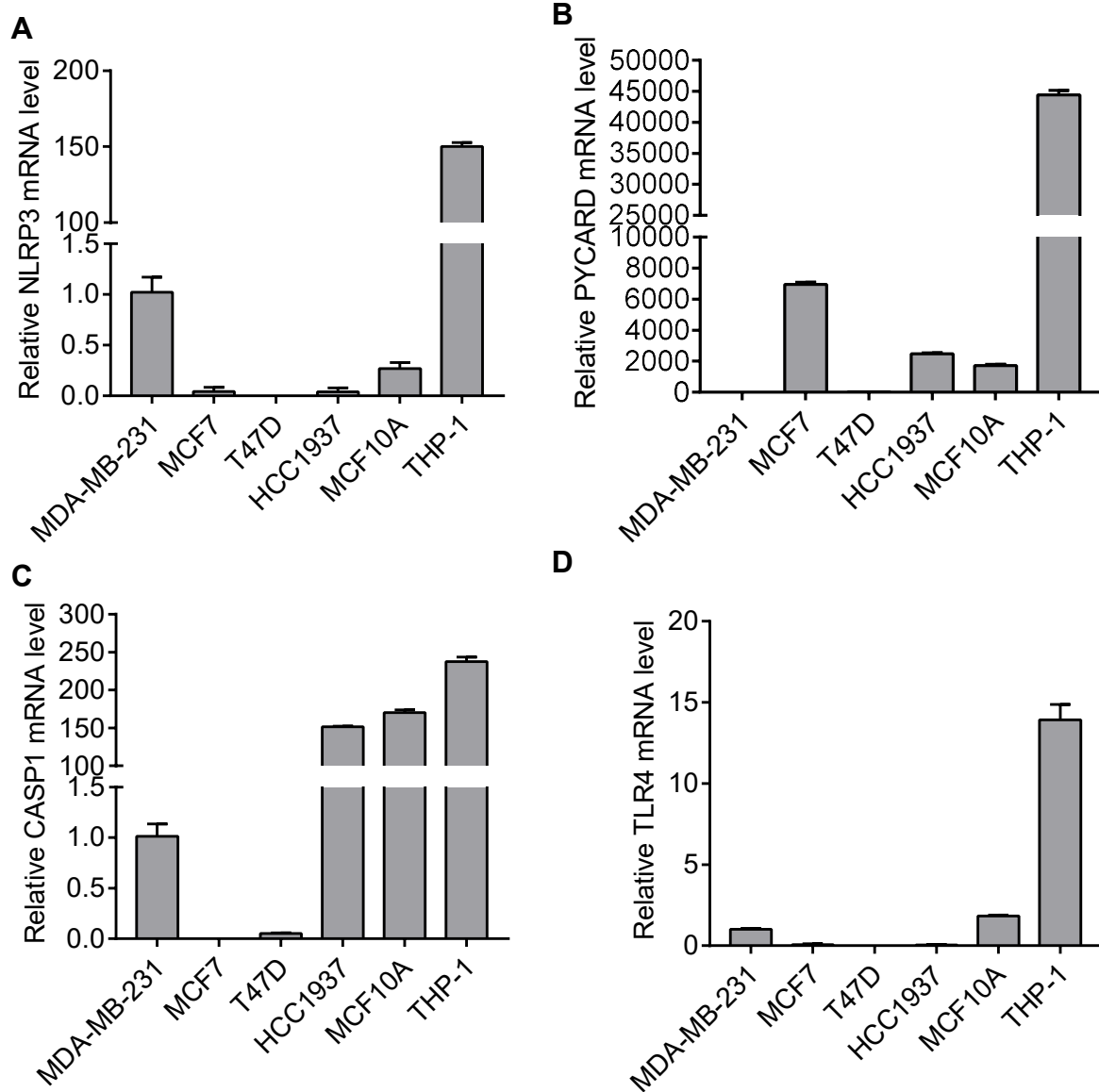
**Figure S5. BRCA1 is required for stress-induced mitochondrial fission via mediating AMPK activation.** **A)** Immunoblot analysis of Flag-BRCA1 in Flag-tagged BRCA1 (Flag-BRCA1) MEFs compared with wildtype (WT) MEFs. **B)** Immunoblot analysis of BRCA1 level on mitochondria in Flag-BRCA1 MEFs under CCCP (30  $\mu$ M) treatment. TOM20 and Lamin A are loading control for mitochondrial and nuclear protein, respectively. **C)** Representative images of BRCA1 in Flag-BRCA1 MEFs stained with Flag antibody. The cells were treated with or not CCCP. Scale bar, 50  $\mu$ m. **D)** BRCA1 is required for Oligomycin-induced AMPK activation in 293T cells. shCtrl and shBRCA1 293T cells were treated by Oligomycin (10  $\mu$ M) for indicated time points. **E)** BRCA1 has no effect on AICAR-induced AMPK activation. *Brca1*<sup>flox/flox</sup> MEFs infected with AdGFP or AdCre were treated with AICAR (2.5 mM) for 1 h. **F, G)** BRCA1 deficiency impairs CCCP-induced mitochondrial fission. (F) The BRCA1 levels in shCtrl and shBRCA1 HeLa-HA-Parkin; (G) Mitochondrial morphology of shCtrl and shBRCA1 HeLa-HA-Parkin with or without CCCP treatment for indicated time points. Mitochondria were marked by MitoDsRed. Arrows indicate mitochondrial morphology. **H)** Loss of DRP1 blocks CCCP-induced mitophagy. shCtrl and shDRP1 HeLa-mCherryParkin were treated with CCCP for indicated time points. Data are representative of at least two independent experiments.



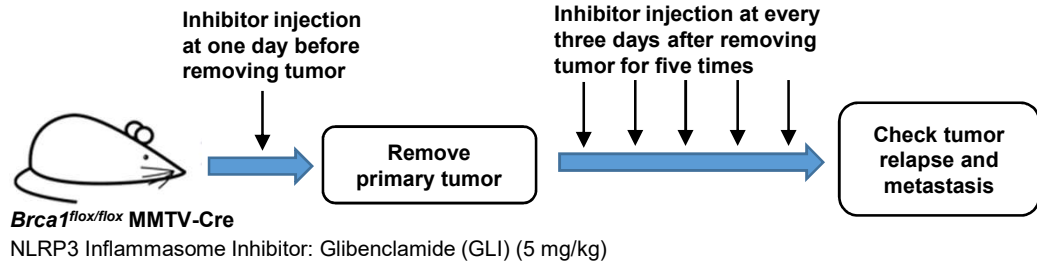
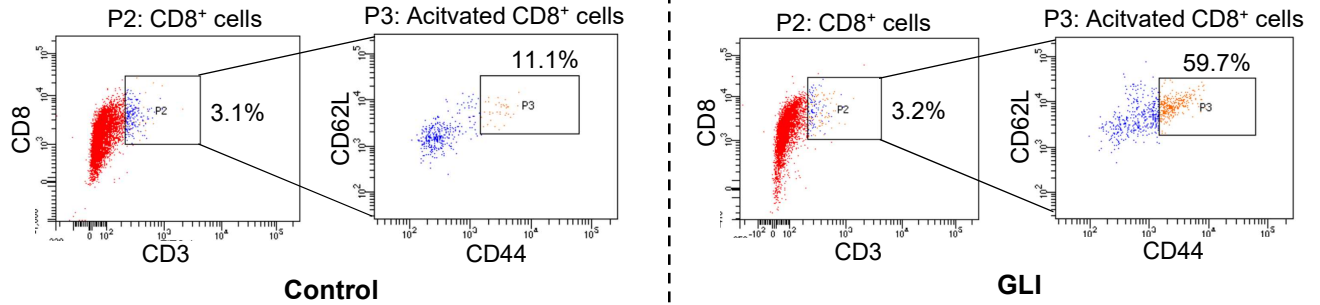
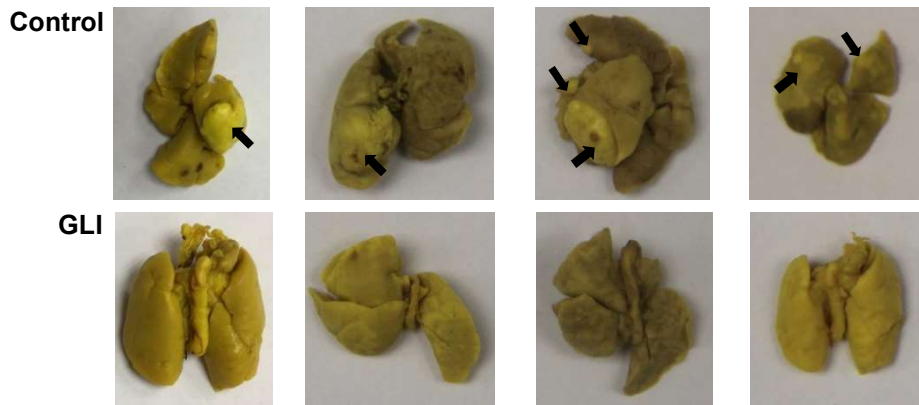
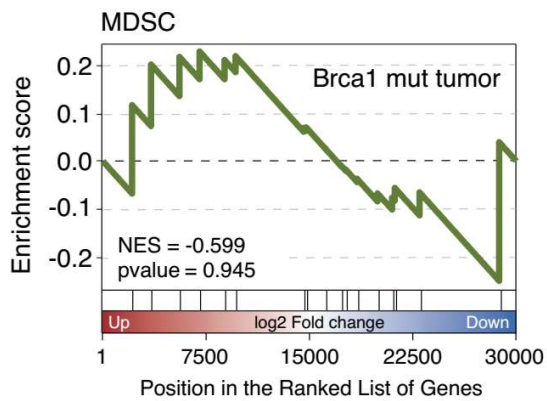
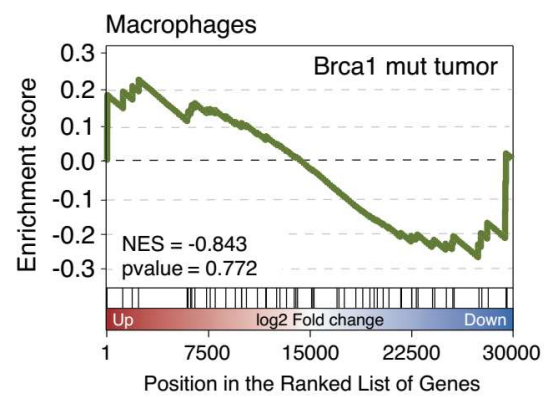
**Figure S6. CCCP-induced AMPK activation is independent on LKB1 or CaMMKβ.** **A)** LKB1 KD has no effect on CCCP-induced AMPK activation. shCtrl and shLKB1 293T cells were treated by CCCP for indicated time points. **B)** CaMMKβ KD has no effect on CCCP-induced AMPK activation. shCtrl and shCaMMKβ 293T cells were treated by CCCP for indicated time points. **C)** Immunoblot analysis of AMPK activation in shCtrl and shBRCA1 Hela cells that lack of LKB1. Data are representative of at least two independent experiments.



**Figure S7. Inflammatory response significantly increases in *Brca1* mutant mammary gland.** GSEA plot of enrichment in “inflammatory response” gene set upregulated in *Brca1* MT mammary glands.



**Figure S8. The expression of inflammasome associated genes in different cell lines. A) NLRP3 mRNA levels. B) PYCARD mRNA levels. C) CASP1 mRNA levels. D) TLR4 mRNA levels.**

**A****B****C****D****E**



**Figure S9. Inhibition of inflammasome activation attenuates *Brca1* mutant mammary tumor recurrence and metastasis.** **A)** The flowchart of animal experiments. *Brca1<sup>flox/flox</sup>* MMTV-Cre mice developing mammary tumors (diameter around 1 cm) were i.p. injected with PBS or NLRP3 inflammasome inhibitor Glibenclamide (GLI, 5 mg/kg) one day before primary tumor removal. Then, the mice were administrated with PBS or GLI by i.p. continuous injection every three days for five times. **B)** Flow cytometric gating strategy for CD8<sup>+</sup> T cell analysis. **C)** Representative images of lung metastasis of tumors after primary tumor removal. Arrows indicate metastatic tumors on lung. **D)** GSEA plot of enrichment in “MDSC” gene set has no significant difference in B.T versus Trp53.T. **E)** GSEA plot of enrichment in “Macrophages” gene set has no significant difference in B.T versus Trp53.T.